AMENDMENTS TO THE CLAIMS

Claims 1 to 25 (previously cancelled)

Claim 26 (currently amended)

A single stranded oligonucleotide OY emprising having 9 to 42 nucleotides of the sequence 9 to 42 nucleotides of the sequence $Y_1.Y_2-Y_3-Y_4-Y_5$ wherein Y_1 is a nucleotide sequence of 1 to 12 nucleotides or is absent suppressed, Y_2 is a trinucleotide which encodes for Gly, Y_3 is a nucleotide coding for Arg or Lys, Y_4 is a nucleotide coding for Arg or Lys and Y_5 is a nucleotide sequence, $Y_6-Y_7-Y_8-Y_9$ wherein Y_6 is a trinucleotide which codes for Ser, Thr or Tyr, Y_7 is a trinucleotide which codes for any amino acid, Y_8 is a trinucleotide which codes for Glu or Asp and Y_9 is a nucleotide sequence of 1 to 12 nucleotides or Y_5 is absent suppressed with the exception of CGACACUCCA CCAUA.

Claim 27 (currently amended)

A oligonucleotide of claim 26 wherein Y_1 and Y_9 are absent suppressed.

Claim 28 (previously added)

An oligonucleotide of claim 27 wherein Y_2 is a trinucleotide which codes for Gly, Y_3 is a trinucleotide which codes for Lys, Y_4 is a trinucleotide which codes for Arg and Y_5 is a sequence of 3 trinucleotides which code for Ser-Ala-glu.

Claim 29 (currently amended)

A single-stranded oligonucleotide OZ eomprising having 15 to 39 nucleotides and hybridizes under mild or strigent conditions with a consensus signal characteristic of amidated polypeptide hormones with the sequence having the formula

$$Z_1-Z_2-Z_3-Z_4-Z_5-Z_6-Z_7$$

wherein is a nucleotide sequence of 1 to 12 nucleotide or is absent suppressed, Z_2 and Z_3 are two trinucleotides which code for Leu, Z_4 and Z_5 are two trinucleotide which code for any two amino acids, Z_6 is a trinucleotide which codes for Leu and Z_7 is a nucleotide sequence of 1 to 12 nucleotides or its absent suppressed.

Claims 30 to 32 (currently cancelled)